aroundEC2 mm10_dna:1-5873

UTR CNS

Alignment 1 EC2 PEc13_2 (+) 100% 1-4158 Criteria: 70%, 100 bp Regions: 1 Alignment 2 human 50% 100% hg38_dna (+) 119790-125892 Criteria: 70%, 100 bp Regions: 7 Alignment 3 50% 100% sheep oviAri4_dna (+) 134844-140833 Criteria: 70%, 100 bp Regions: 8 JAMAY YA М Alignment 4 50% 100% pig susScr11_dna (+) 131809-137474 Criteria: 70%, 100 bp Regions: 9 \sqrt{N} Alignment 5 50% 100% oppossum monDom5_dna (+) 166443-173475 Criteria: 70%, 100 bp Regions: 5 Alignment 6 50% 100% chicken galGal6_dna (+) 30713-36576 Criteria: 70%, 100 bp Regions: 1 Alignment 7 50% 100% cobra CM019149.1_166319490_166896960 (+) 466357-472226 Criteria: 70%, 100 bp Regions: 0 Alignment 8 50% 100% org9 no alignment Criteria: 70%, 100 bp Regions: 0 X-axis: aroundEC2 Resolution: 7 Window size: 100 bp 0k 1k 2k 3k 4k 5k contig gene exon

aroundEC2 mm10_dna:1-5873

Alignment 1 EC2 PEc13_2 (+) 1-4158 Criteria: 70%, 100 bp Regions: 1 Alignment 2 human hg38_dna (+) 119790-125892 Criteria: 70%, 100 bp Regions: 7 Alignment 3 sheep oviAri4_dna (+) 134844-140833 Criteria: 70%, 100 bp Regions: 8 Alignment 4 pig susScr11_dna (+) 131809-137474 Criteria: 70%, 100 bp Regions: 9 Alignment 5 oppossum monDom5_dna (+) 166443-173475 Criteria: 70%, 100 bp Regions: 5 Alignment 6 chicken galGal6_dna (+) 30713-36576 Criteria: 70%, 100 bp Regions: 1 Alignment 7 cobra CM019149.1_166319490_166896960 (+) 466357-472226 Criteria: 70%, 100 bp Regions: 0 Alignment 8 org9 no alignment Criteria: 70%, 100 bp Regions: 0 X-axis: aroundEC2 Resolution: 7 Window size: 100 bp contig gene exon

UTR

